



SEQUENCE LISTING

<110> Abe, Hiroko
Shimma, Yoh-ichi
Jigami, Yoshifumi

<120> NUCLEIC ACIDS, EXPRESSION VECTORS AND
HOST CELLS FOR MAKING CHIMERIC NUCLEIC ACIDS AND METHODS FOR
PRODUCING IMMOBILIZED POLYPEPTIDES

<130> 13558-004001

<140> 09/989,975

<141> 2001-11-21

<150> JP 2001-190524

<151> 2001-06-22

<150> JP 2000-354396

<151> 2000-11-21

<160> 14

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 341

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 1

Met	Gln	Tyr	Lys	Lys	Ser	Leu	Val	Ala	Ser	Ala	Leu	Val	Ala	Thr	Ser
1				5					10					15	
Leu	Ala	Ala	Tyr	Ala	Pro	Lys	Asp	Pro	Trp	Ser	Thr	Leu	Thr	Pro	Ser
			20					25					30		
Ala	Thr	Tyr	Lys	Gly	Gly	Ile	Thr	Asp	Tyr	Ser	Ser	Thr	Phe	Gly	Ile
		35				40						45			
Ala	Val	Glu	Pro	Ile	Ala	Thr	Thr	Ala	Ser	Ser	Lys	Ala	Lys	Arg	Ala
	50					55					60				
Ala	Ala	Ile	Ser	Gln	Ile	Gly	Asp	Gly	Gln	Ile	Gln	Ala	Thr	Thr	Lys
65				70				75						80	
Thr	Thr	Ala	Ala	Ala	Val	Ser	Gln	Ile	Gly	Asp	Gly	Gln	Ile	Gln	Ala
			85					90						95	
Thr	Thr	Lys	Thr	Lys	Ala	Ala	Ala	Val	Ser	Gln	Ile	Gly	Asp	Gly	Gln
		100						105						110	
Ile	Gln	Ala	Thr	Thr	Lys	Thr	Thr	Ser	Ala	Lys	Thr	Thr	Ala	Ala	Ala
	115					120							125		
Val	Ser	Gln	Ile	Gly	Asp	Gly	Gln	Ile	Gln	Ala	Thr	Thr	Lys	Thr	Lys
	130					135					140				
Ala	Ala	Ala	Val	Ser	Gln	Ile	Gly	Asp	Gly	Gln	Ile	Gln	Ala	Thr	Thr
145					150					155					160
Lys	Thr	Thr	Ala	Ala	Ala	Val	Ser	Gln	Ile	Gly	Asp	Gly	Gln	Ile	Gln
			165					170						175	
Ala	Thr	Thr	Lys	Thr	Thr	Ala	Ala	Ala	Val	Ser	Gln	Ile	Gly	Asp	Gly
			180					185						190	

Gln Ile Gln Ala Thr Thr Asn Thr Thr Val Ala Pro Val Ser Gln Ile
 195 200 205
 Thr Asp Gly Gln Ile Gln Ala Thr Thr Leu Thr Ser Ala Thr Ile Ile
 210 215 220
 Pro Ser Pro Ala Pro Ala Pro Ile Thr Asn Gly Thr Asp Pro Val Thr
 225 230 235 240
 Ala Glu Thr Cys Lys Ser Ser Gly Thr Leu Glu Met Asn Leu Lys Gly
 245 250 255
 Gly Ile Leu Thr Asp Gly Lys Gly Arg Ile Gly Ser Ile Val Ala Asn
 260 265 270
 Arg Gln Phe Gln Phe Asp Gly Pro Pro Pro Gln Ala Gly Ala Ile Tyr
 275 280 285
 Ala Ala Gly Trp Ser Ile Thr Pro Glu Gly Asn Leu Ala Ile Gly Asp
 290 295 300
 Gln Asp Thr Phe Tyr Gln Cys Leu Ser Gly Asn Phe Tyr Asn Leu Tyr
 305 310 315 320
 Asp Glu His Ile Gly Thr Gln Cys Asn Ala Val His Leu Gln Ala Ile
 325 330 335
 Asp Leu Leu Asn Cys
 340

<210> 2

<211> 413

<212> PRT

<213> *Saccharomyces cerevisiae*

<400> 2

Met Gln Tyr Lys Lys Thr Leu Val Ala Ser Ala Leu Ala Ala Thr Thr
 1 5 10 15
 Leu Ala Ala Tyr Ala Pro Ser Glu Pro Trp Ser Thr Leu Thr Pro Thr
 20 25 30
 Ala Thr Tyr Ser Gly Gly Val Thr Asp Tyr Ala Ser Thr Phe Gly Ile
 35 40 45
 Ala Val Gln Pro Ile Ser Thr Thr Ser Ser Ala Ser Ser Ala Ala Thr
 50 55 60
 Thr Ala Ser Ser Lys Ala Lys Arg Ala Ala Ser Gln Ile Gly Asp Gly
 65 70 75 80
 Gln Val Gln Ala Ala Thr Thr Thr Ala Ser Val Ser Thr Lys Ser Thr
 85 90 95
 Ala Ala Ala Val Ser Gln Ile Gly Asp Gly Gln Ile Gln Ala Thr Thr
 100 105 110
 Lys Thr Thr Ala Ala Ala Val Ser Gln Ile Gly Asp Gly Gln Ile Gln
 115 120 125
 Ala Thr Thr Lys Thr Thr Ser Ala Lys Thr Thr Ala Ala Ala Val Ser
 130 135 140
 Gln Ile Ser Asp Gly Gln Ile Gln Ala Thr Thr Thr Leu Ala Pro
 145 150 155 160
 Lys Ser Thr Ala Ala Val Ser Gln Ile Gly Asp Gly Gln Val Gln
 165 170 175
 Ala Thr Thr Thr Leu Ala Pro Lys Ser Thr Ala Ala Ala Val Ser
 180 185 190
 Gln Ile Gly Asp Gly Gln Val Gln Ala Thr Thr Lys Thr Thr Ala Ala
 195 200 205
 Ala Val Ser Gln Ile Gly Asp Gly Gln Val Gln Ala Thr Thr Lys Thr
 210 215 220
 Thr Ala Ala Ala Val Ser Gln Ile Gly Asp Gly Gln Val Gln Ala Thr
 225 230 235 240

Thr	Lys	Thr	Thr	Ala	Ala	Ala	Val	Ser	Gln	Ile	Gly	Asp	Gly	Gln	Val
				245					250					255	
Gln	Ala	Thr	Thr	Lys	Thr	Thr	Ala	Ala	Ala	Val	Ser	Gln	Ile	Thr	Asp
				260				265					270		
Gly	Gln	Val	Gln	Ala	Thr	Thr	Lys	Thr	Thr	Gln	Ala	Ala	Ser	Gln	Val
				275			280					285			
Ser	Asp	Gly	Gln	Val	Gln	Ala	Thr	Thr	Ala	Thr	Ser	Ala	Ser	Ala	Ala
	290					295					300				
Ala	Thr	Ser	Thr	Asp	Pro	Val	Asp	Ala	Val	Ser	Cys	Lys	Thr	Ser	Gly
305				310						315					320
Thr	Leu	Glu	Met	Asn	Leu	Lys	Gly	Gly	Ile	Leu	Thr	Asp	Gly	Lys	Gly
				325					330					335	
Arg	Ile	Gly	Ser	Ile	Val	Ala	Asn	Arg	Gln	Phe	Gln	Phe	Asp	Gly	Pro
			340				345					350			
Pro	Pro	Gln	Ala	Gly	Ala	Ile	Tyr	Ala	Ala	Gly	Trp	Ser	Ile	Thr	Pro
		355					360					365			
Asp	Gly	Asn	Leu	Ala	Ile	Gly	Asp	Asn	Asp	Val	Phe	Tyr	Gln	Cys	Leu
	370					375					380				
Ser	Gly	Thr	Phe	Tyr	Asn	Leu	Tyr	Asp	Glu	His	Ile	Gly	Ser	Gln	Cys
385				390						395					400
Thr	Pro	Val	His	Leu	Glu	Ala	Ile	Asp	Leu	Ile	Asp	Cys			
				405					410						

```
<210> 3
<211> 44
<212> DNA
<213> Artificial Sequence
```

<220>
<223> Synthetic primer

44

```
<210> 4
<211> 31
<212> DNA
<213> Artificial Sequence
```

<220>
<223> Synthetic primer

31

```
<210> 5
<211> 35
<212> DNA
<213> Artificial Sequence
```

<220>
<223> Synthetic primer

35

<210> 6

<211> 47
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic primer

<400> 6
 gggggcccggg ctaggatgat gggtttcaaaa gattttgaat atgatcc 47

<210> 7
 <211> 35
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic primer

<400> 7
 cccgtcgaca atcctatctg cgtgtgtctc aagac 35

<210> 8
 <211> 35
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic primer

<400> 8
 cccctcgagt caggtgaacc aagccgctat gccgc 35

<210> 9
 <211> 39
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic primer

<400> 9
 gggggggtcga cagcaatata ttccgagttc catctccgc 39

<210> 10
 <211> 42
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Synthetic primer

<400> 10
 ggggggtcga gctactcacg gaattttttc cagttttttg gc 42

<210> 11
 <211> 38
 <212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer

<400> 11

gggggggagct catgcaatac aaaaagactt tggttgcc

38

<210> 12

<211> 68

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer

<400> 12

cccccgcggc cgcttgtca tcgtcatcct tgtagtcaca gtctatcaaa tcgatatgctt
ccaagtgg

60

68

<210> 13

<211> 49

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer

<400> 13

ggggggcggc cgcaaatgat gcgcttatac gatcaagcaa tgtaaacag

49

<210> 14

<211> 38

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic primer

<400> 14

gggggcccgg gctagctttg ttcgtgtcta gaattttc

38